

FIG. 1B

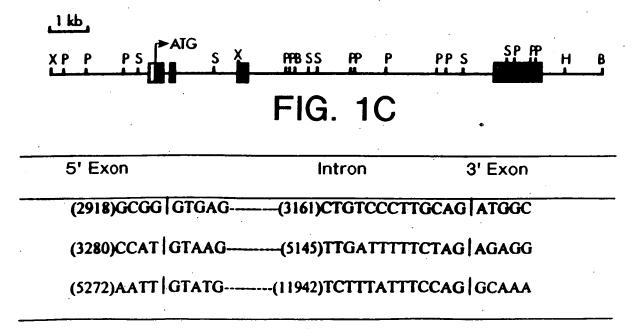
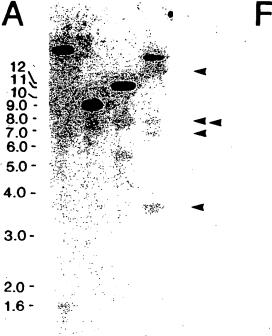


FIG. 1D

FIG. 2A

Ecopi Barnill Dal Hindli

FIG. 2B



450 -375 -295 -225 -

23-

FIG. 2C

23.9 -

9.4 -

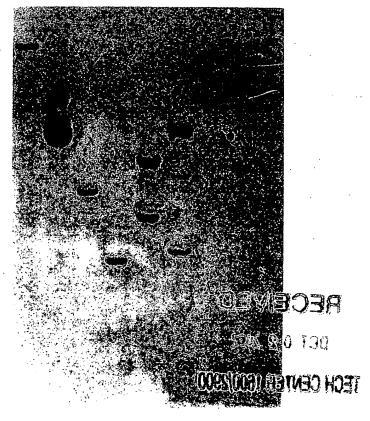
6.7 -

4.4 -

2.3 -

2.0-

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9.5 -7.5 -

9.5 **-** 7.5 **-**

4.4 -

4.4 -

2.4 -

2.4 -

1.35 _

1.35 -

FIG. 3A

FIG. 3B

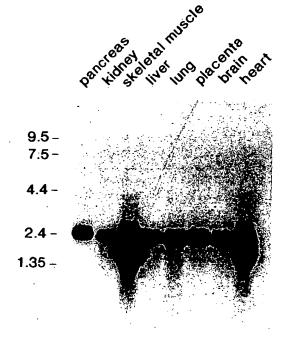


FIG. 3C

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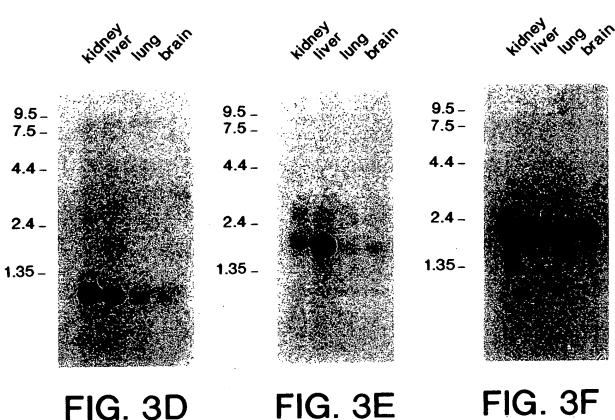


FIG. 3E FIG. 3D

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FIG. 4B-2

		٠.										ì
Majority	0	L49433.PRO L49441.PRO	9	P41437.PRO	U19251.PRO	• ·	U32974.PRO	U36842.PRO	U45878.PRO	U45879. PRO	U45880.PRO	U45881.PRO
GOEYUS	400	GOEFND	GRUM	GRDFW-	SSAEVII	GPSYVIS	GOEYIN	GOEYIN	GOEFIR	GOEFTID	GOEYIN	(GPAVVIS
CEFLLLAK	390	COEVILAR COEVILAR	SAYVOLK	GEYVILIVA	CEFTONM	公包包工工具	公KY打可受	公KY打り 県	DEVATIRITY	CEETTENTS	AKYALIED	GOEVULUA AOEVULUA
PWEEHAKWFPR	380	PWIEDJAKWE PR	WEDDENEWADE	CANO COCHA ELMY DE	HELDDATERCE FOL	PWEIDTAKWSEK	EWECTAKWAYEG	PWEQHAKWYPG	PWV ODAKWE/PR	CANDIDAKWENE	EWELOGIA KWIYING	DWEETA KWSOK
RDWEPGDD	370	RCWEPGDO RSWOKEDE	KLWE PECO	KUWEPDAA	丘氏がの丘合かり	RSWOKEDE	FLOOKENS BO	TD WICHS ED	RCM回SCCC	RCM可SCCC	TOWKIPSED	BISKAKABB
YVGIGDKVKCFHCDGGLRDWEPGDDPWEEHAKWFPRCEFLLLAKGQEYVS	360	YNDRNDDVKCFCCDGGL YOKTGDOVRCFHCNIGL	V TGY GON TWO FY CDGGL	文化公のGDKTRACCCTAGA	1/1/61/KD1/1/0CF/SCGGG1	MONTGOONE CONTONICA	AIDEDDDDDDDDDDDDDDDDDDDDDDDDDDDDDDDDDDD	ALIZEGIAN SKOLLYGGEN	NVON SIZED SKY MCCF TO CONTROLL TO CONTROL	MY CRIMINATION OF CHARACTER CONTROLL OF CHARACTER CONTROLL OF CHARACTER CONTROL OF CONTROL OF CONTROL OF CHARACTER CONTROL OF CHARACTER CONTROL OF CONTROL	ALGEBANDA CALACAGA	Montiedolarchied

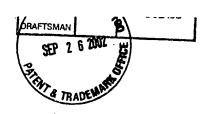


FIG. 4A

10 30 40

IIIIII 10 40

MGAPTLPPAW OPFLKDHRIS TFKNWPFLEG CACTPERMAE 40

AGFIHCPTEN EPDLAOCFFC FKELEGWEPD DDPIEEHKKH 80

SSGCAFLSVK KOFEELTLGE FLKLDRERAK NKIAKETNNK 120

KKEFEETAKK VRRAIEOLAA MD 142

FIG. 4B-1

-TPQELAKAGFYYLGRGDQVQCFACGGKLA Majority EEARLVTFONWPD-AFI

	RO	RO	RO	RO	RO.	PRO	RO .	RO	RO .	PRO	RO	PRO	I. PRO
	33.	L49441.F	P41436.E	•	U19251.F	U32373.E	2974.	•	φ.	<u>ი</u>	U45880.F	H	SURVIVIN
250	ACCOMIS	MCNOOVIA MCNOOVIA	FOKVEIM	FOKVEIT	5(56651)6	MONOTONIA		いが存在所	A A A A A A A A A A A A A A A A A A A	ACCORDS	公公公 公公公 公公 公公 公 公 公 公 公 公 公 之 分 、 、 、 、 、 、 、 、 、 、 、 、 、	MONGAV TA	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
240	GURNAGE	TATAKON	SDEVECA	GODONACA.	90年/06年	TOTAL TOTAL	GHOVOCE	DOVOCE	GARZESTE STATES	GEN SACE	回 日 日 日 日 日 日 日 日 日 日 日 日 日	TATAKOV.	
230	可,ARAG的VY区GP	ALLA KAGGEYY LINIR	TMAKNGEYYLGR	RMAASGEVYLGR	-1,556,65,VEUGK	ALLA KÁGÉTYYLINIR	bíjaskárvytár	DJASAGUYYEGA	DJARAGEY YIGP	ELJARÁGEIY VIJGP	ELFASKGLYVEGT	ALLAKAGEVYLINE	RM <u>A度為</u> 住戶II
220	1SPA	IIII	工SPE	LEPS	G18PCV	$\overline{\text{II}}$ – – – $\overline{\text{TPQ}}$	LTPR	11TPR	LSPT	LSPS	LTPR	DalI	G-CACT PE
210	7 BEASPELITY SMOSSINGELISE	HANBIN TEKDWEN-PI	回知が打いて回じばがらV-SF	KAAKUGIIYIMWBW-OF	9 EEARTASERNWFEYVO		3 PRINCESCONNICTION	3 FRANKERONWEDYALI	9 END STITE TEND TWO LITE	4 EFARELDY HIMM P-LTE	A. DEAKTESTON/ADVAIN	DANKINTEKUMEN-PN	XDHKITSEEKKWが近-LE
	17			ω _.	. Т	 1 '	10	16	16	رين ص	9	다 (단 ()	

MWx10⁻³

200 -

97.4 -

69 -

46 -

30 -

21.5 -14.3 -

Survivin.

FIG. 4C-1

MWx10-3

200 -

97.4 -

69 -

46 -

30-

21.5-

14.3-



RblgG

FIG. 4C-2

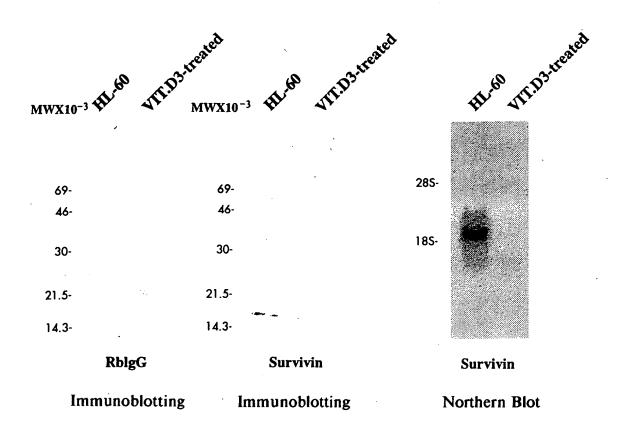


FIG. 5A FIG. 5B FIG. 5C



FIG. 6A



FIG. 6C

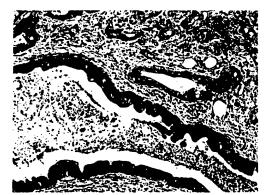


FIG. 6E

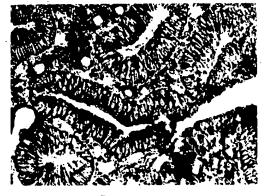


FIG. 6G

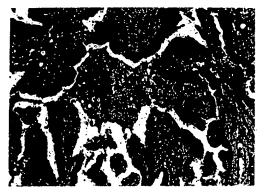


FIG. 6B

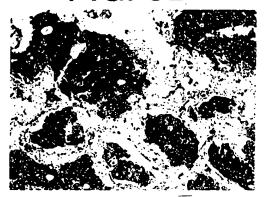


FIG. 6D

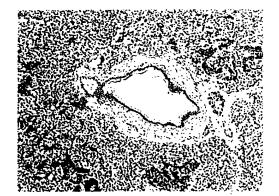


FIG. 6F



FIG. 6H

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MW x10⁻³

Jecto

Antisense

200 -

97.4 -

69 -

46 -

30 -

21.5 -

14.3 -

FIG. 7A

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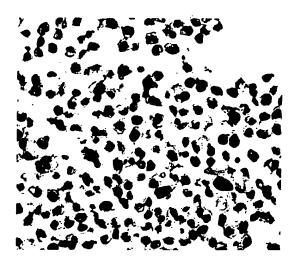


FIG. 7B-1

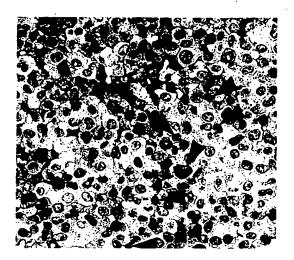


FIG. 7B-2

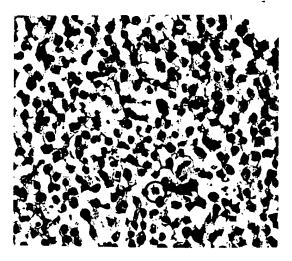


FIG. 7B-3

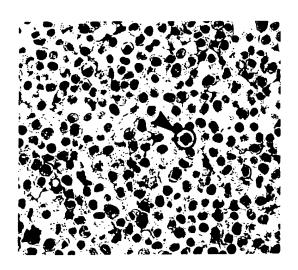


FIG. 7B-4

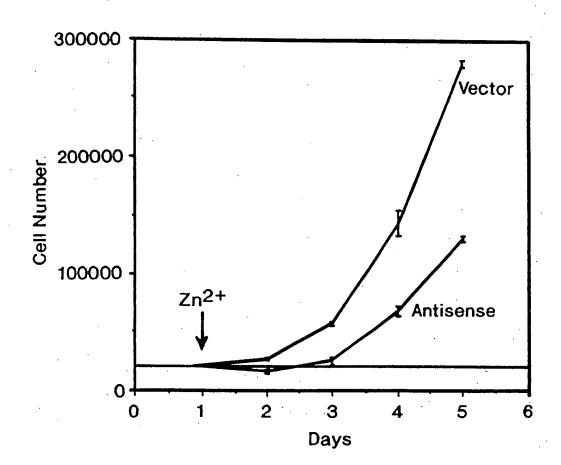


MARC IM

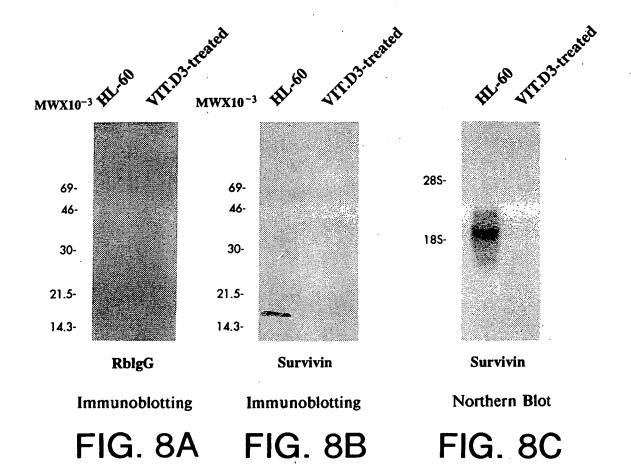




FIG. 7C



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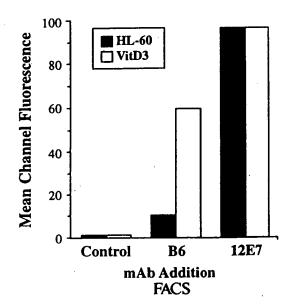


FIG. 8D

FIG. 9

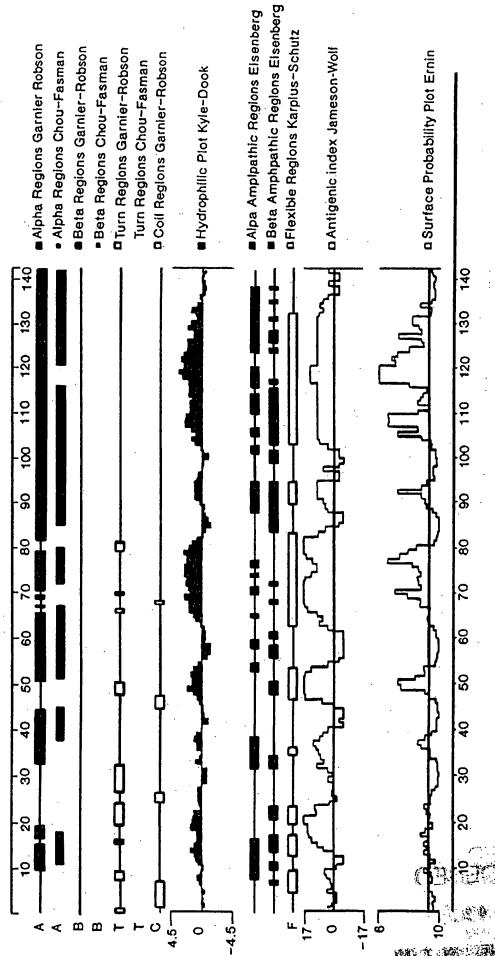




FIG. 10A

translation=MGAPTLPPAWQPFLKDHRISTFKNWPFLEGCACTPERMAEAGFIHCP TENEPDLAQCFFCFKELEGWEPDDDPIEEHKKHSSGCAFLSVKKQFEELTLGEFL KLDRERAKNKIAKETNNKKKEFEETAKKVRRAIEQLAAMD

			•		
1 .	TCTAGACATG	CGGATATATT	CAAGCTGGGC	ACAGCACAGC	AGCCCCACCC
51	CAGGCAGCTT	GAAATCAGAG	CTGGGGTCCA	AAGGGACCAC	ACCCCGAGGG
101	ACTGTGTGGG	GGTCGGGGCA	CACAGGCCAC	TGCTTCCCCC	CGTCTTTCTC
151	AGCCATTCCT	GAAGTCAGCC	TCACTCTGCT	TCTCAGGGAT	TTCAAATGTG
201	CAGAGACTCT	GGCACTTTTG	TAGAAGCCCC	TTCTGGTCCT	AACTTACACC
251	TGGATGCTGT	GGGGCTGCAG	CTGCTGCTCG	GGCTCGGGAG	GATGCTGGGG
301	GCCCGGTGCC	CATGAGCTTT	TGAAGCTCCT	GGAACTCGGT	TTTGAGGGTG
351	TTCAGGTCCA	GGTGGACACC	TGGGCTGTCC	TTGTCCATGC	ATTTGATGAC
401	ATTGTGTGCA	GAAGTGAAAA	GGAGTTAGGC	CGGGCATGCT	GGCTTATGCC
451	TGTAATCCCA	GCACTTTGGG	AGGCTGAGGC	GGGTGGATCA	CGAGGTCAGG
501	AGTTCAATAC	CAGCCTGGCC	AAGATGGTGA	AACCCCGTCT	CTACTAAAAA
551	TACAAAAAAA	TTAGCCGGGC	ATGGTGGCGG	GCGCATGTAA	TCCCAGCTAC
601	TGGGGGGGCT	GAGGCAGAGA	ATTGCTGGAA	CCCAGGAGAT	GGAGGTTGCA
651	GTGAGCCAAG	ATTGTGCCAC	TGCACTGCAC	TCCAGCCTGG	CGACAGAGCA
701	AGACTCTGTC	TCAAAAAAAA	AAAAAAAAG	TGAAAAGGAG	TTGTTCCTTT
751	CCTCCCTCCT	GAGGGCAGGC	AACTGCTGCG	GTTGCCAGTG	GAGGTGGTGC
801	GTCCTTGGTC	TGTGCCTGGG	GGCCACCCCA	GCAGAGGCCA	TGGTGGTGCC
851	AGGGCCCGGT	TAGCGAGCCA	ATCAGCAGGA	CCCAGGGGCG	ACCTGCCAAA
901	GTCAACTGGA	TTTGATAACT	GCAGCGAAGT	TAAGTTTCCT	GATTTTGATG
951	ATTGTGTTGT	GGTTGTGTAA	GAGAATGAAG	TATTTCGGGG	TAGTATGGTA
1001	ATGCCTTCAA	CTTACAAACG	GTTCAGGTAA	ACCACCCATA	TACATACATA
1051	TACATGCATG	TGATATATAC	ACATACAGGG	ATGTGTGTGT	GTTCACATAT
1101	ATGAGGGGAG	AGAGACTAGG	GGAGAGAAAG	TAGGTTGGGG	AGAGGGAGAG
1151	AGAAAGGAAA	ACAGGAGACA	GAGAGAGAGC	GGGGAGTAGA	GAGAGGGAAG
1201	GGGTAAGAGA	GGGAGAGGAG	GAGAGAAAGG	GAGGAAGAAG	CAGAGAGTGA
1251	ATGTTAAAGG	AAACAGGCAA	AACATAAACA	GAAAATCTGG	GTGAAGGGTA
1301	TATGAGTATT	CTTTGTACTA	TTCTTGCAAT	TATCTTTTAT	TTAAATTGAC
1351	ATCGGGCCGG	GCGCAGTGGC	TCACATCTGT	AATCCCAGCA	CTTTGGGAGG
1401	CCGAGGCAGG	CAGATCACTT	GAGGTCAGGA	GTTTGAGACC	AGCCTGGCAA
1451	ACATGGTGAA	ACCCCATCTC	TACTAAAAAT	ACAAAAATTA	GCCTGGTGTG
1501	GTGGTGCATG	CCTTTAATCT	CAGCTACTCG	GGAGGCTGAG	GCAGGAGAAT
1551	CGCTTGAACC	CGTGGCGGGG	AGGAGGTTGC	AGTGAGCTGA	GATCATGCCA
1601	CTGCACTCCA	GCCTGGGCGA	TAGAGCGAGA	CTCAGTTTCA	AATAAATAAA
1651	TAAACATCAA	AATAAAAAGT	TACTGTATTA	AAGAATGGGG	GCGGGGTGGG
1701	AGGGGTGGGG	AGAGGTTGCA	ATAAATA	AATAAATAA	TAAACCCCAA
1751	AATGAAAAAG	ACAGTGGAGG	CACCAGGCCT	GCGTGGGGCT	GGAGGGCTAA
1801	TAAGGCCAGG	CCTCTTATCT	CTGGCCATAG	AACCAGAGAA	GTGAGTGGAT
1851	GTGATGCCCA	GCTCCAGAAG	TGACTCCAGA	ACACCCTGTT	CCAAAGCAGA
1901	GGACACACTG	ATTTTTTTTT	TAATAGGCTG	CAGGACTTAC	TGTTGGTGGG
1951	ACGCCCTGCT	TTGCGAAGGG	AAAGGAGGAG	TTTGCCCTGA	GCACAGGCCC
2001	CCACCCTCCA	CTGGGCTTTC	CCCAGCTCCC	TTGTCTTCTT	ATCACGGTAG
2051	TGGCCCAGTC	CCTGGCCCCT	GACTCCAGAA	GGTGGCCCTC	CTGGAAACCC
2101	AGGTCGTGCA	GTCAACGATG	TACTCGCCGG	GACAGCGATG	TCTGCTGCAC
2151	TCCATCCCTC	CCCTGTTCAT	TTGTCCTTCA	TGCCCGTCTG	GAGTAGATGC

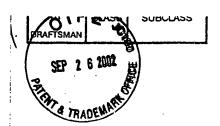


FIG. 10B

2201	TTTTTGCAGA	GGTGGCACCC	TGTAAAGCTC	TCCTGTCTGA	CTTTTTTTTT
2251				TAGGCTGGAG	•
2301	CAATCTCAGC			CGGGTTCAAG	
2351	GCCTCAGCCT			GGCATGCACC	-
2401				GTTTCACCGT	
2451	•			CTCCTGCCTA	
2501				ACCCGGCCTG	
2551		•		GGCAGGGACG	
2601				GAGCCACGCG	
2651			4	GCCTCTACTC	
2701		•		GCTCCCGACA	•
2751	CGCGCCATTA	ACCGCCAGAT	TTGAATCGCG	GGACCCGTTG	GCAGAGGTGG
			•		
2801	CGGCGGCGGC	•		CCCTGCCTGG	CAGCCCTTTC
2851		•		ACTGGCCCTT	
2901				CCCGGCCTCC	
2951		TTGCCCTGTC			TGGGCCTCGG
3001				GTCCCCAGCG	AGGCCACTGT
3051				CCCTCCCTGC	
3101				CCGGGCTGCC	•
3151				AGGCTGGCTT	
3201	CCCACTGAGA	ACGAGCCAGA	CTTGGCCCAG	TGTTTCTTCT	GCTTCAAGGA
3251	GCTGGAAGGC	TGGGAGCCAG	ATGACGACCC	CATGTAAGTC	TTCTCTGGCC
3301		GGCTTTGTTT	•		TTTGAGTTGC
3351	AAAGACACTT	AGTATGGGAG	GGTTGCTTTC	CACCCTCATT	GCTTCTTAAA
3401	CAGCTGTTGT	GAACGGATAC	CTCTCTATAT	GCTGGTGCCT	TGGTGATGCT
3451	TACAACCTAA	TTAAATCTCA	TTTGACCAAA	ATGCCTTGGG	GTGGACGTAA
3501	GATGCCTGAT	GCCTTTCATG	TTCAACAGAA	TACATCAGCA	GACCCTGTTG
3551	TTGTGAACTC	CCAGGAATGT	CCAAGTGCTT	TTTTTGAGAT	AAAAATTTTT
3601	ACAGTTTAAT	TGAAATATAA	CCTACACAGC	ACAAAAATTA	CCCTTTGAAA
3651	GTGTGCACTT	CACACTTTCG	GAGGCTGAGG	CGGGCGGATC	ACCTGAGGTC
3701	AGGAGTTCAA	GACCTGCCTG	GCCAACTTGG	CGAAACCCCG	TCTCTACTAA
3751	AAATACAAAA	ATTAGCCGGG	CATGGTAGCG	CACGCCCGTA	ATCCCAGCTA
.3801	CTCGGGAGGC	TAAGGCAGGA	GAATCGCTTG	AACCTGGGAG	GCGGAGGTTG
3851	CAGTGAGCCG	AGATTGTGCC	AATGCACTCC	AGCCTCGGCG	ACAGAGCGAG
3901	ACTCCGTCAT	AAAATAAAA	AATTGAAAAA	АААААААGAA	AGAAAGCATA
3951	TACTTCAGTG	TTGTTCTGGA	TTTTTTTCTT	CAAGATGCCT	AGTTAATGAC
4001	AATGAAATTC	TGTACTCGGA	TGGTATCTGT	CTTTCCACAC	TGTAATGCCA
4051	TATTCTTTTC	TCACCTTTTT	TTCTGTCGGA	TTCAGTTGCT	TCCACAGCTT
4101	TAATTTTTTT	CCCCTGGAGA	TCTTAGTATG	TTTGCTATGG	TGGTTATACT
4151	GCATCCCCGT	AATCACTGGG	AAAAGATCAG	TGGTATTCTT	CTTGAAAATG
4201	AATAAGTGTT	ATGATATTTT	CAGATTAGAG	TTACAACTGG	CTGTCTTTTT
4251	GGACTTTGTG	TGGCCATGTT	TTCATTGTAA	TGCAGTTCTG.	GTAACGGTGA
4301	TAGTCAGTTA	TACAGGGAGA	CTCCCCTAGC	AGAAAATGAG	AGTGTGAGCT
4351	AGGGGGTCCC	TTGGGGAACC	CGGGGCAATA	ATGCCCTTCT	CTGCCCTTAA
4401	TCCTTACAGT	GGGCCGGGCA	CGGTGGCTTA	CGCCTGTAAT	ACCAGCACTT
4451				CAGGAGATCG	\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \
4501				AAAATACAAA	
				. 1916	文:"好好,因此,好啊。"

FIG. 10C

4551	GGCGTGGTGG	TGGGCGCCTG	TAGTCCCAGC	TACTCGGGAG	GCTGAGGCAG
4601	GAGAATGGCG	TGAACCCAGG	AGGCGGAGCT	TGCAGTGAGC	CGAGATTGCA
4651	CCACTGCACT	CCAGCCTGGG	CGACAGAATG	AGACTCCGTC	ТСААААААА
4701	AAAAAAAAGA	AAAAAATCTT	TACAGTGGAT	TACATAACAA	TTCCAGTGAA
4751	ATGAAATTAC	TTCAAACAGT	TCCTTGAGAA	TGTTGGAGGG	ATTTGACATG
4801	TAATTCCTTT	GGACATATAC	CATGTAACAC	TTTTCCAACT	AATTGCTAAG
4851	GAAGTCCAGA	TAAAATAGAT	ACATTAGCCA	CACAGATGTG	GGGGGAGATG
4901	TCCACAGGGA	GAGAGAAGGT	GCTAAGAGGT	GCCATATGGG	AATGTGGCTT
4951	GGGCAAAGCA	CTGATGCCAT	CAACTTCAGA	CTTGACGTCT	TACTCCTGAG
5001	GCAGAGCAGG	GTGTGCCTGT	GGAGGGCGTG	GGGAGGTGGC	CCGTGGGGAG
5051	TGGACTGCCG	CTTTAATCCC	TTCAGCTGCC	TTTCCGCTGT	TGTTTTGATT
5101	TTTCTAGAGA	GGAACATAAA	AAGCATTCGT	CCGGTTGCGC	TTTCCTTTCT
5151	GTCAAGAAGC	AGTTTGAAGA	ATTAACCCTT	GGTGAATTTT	TGAAACTGGA
5201	CAGAGAAAGA	GCCAAGAACA	AAATTGTATG	TATTGGGAAT	AAGAACTGCT
5251	CAAACCCTGT	TCAATGTCTT	TAGCACTAAA	CTACCTAGTC	CCTCAAAGGG
5301	ACTCTGTGTT	TTCCTCAGGA	AGCATTTTTT	TTTTTTTCT	GAGATAGAGT
5351	TTCACTCTTG	TTGCCCAGGC	TGGAGTGCAA	TGGTGCAATC	TTGGCTCACT
5401	GCAACCTCTG	CCTCTCGGGT	TCAAGTGATT	CTCCTGCCTC	AGCCTCCCAA
5451	GTAACTGGGA	TTACAGGGAA	GTGCCACCAC	ACCCAGCTAA	TTTTTGTATT
5501	TTTAGTAGAG	ATGGGGTTTC	ACCACATTGC	CCAGGCTGGT	CTTGAACTCC
5551	TGACCTCGTG	ATTCGCCCAC	CTTGGCCTCC	CAAAGTGCTG	GGATTACAGG
5601	CGTGAACCAC	CACGCCTGGC	TTTTTTTTT	TTGTTCTGAG	ACACAGTTTC
5651	ACTCTGTTAC	CCAGGCTGGA	GTAGGGTGGC	CTGATCTCGG	ATCACTGCAA
5701	CCTCCGCCTC	CTGGGCTCAA	GTGATTTGCC	TGCTTCAGCC	TCCCAAGTAG
5 7 51	CCGAGATTAC	AGGCATGTGC	CACCACACCC	AGGTAATTTT	TGTATTTTTG
5801	GTAGAGACGA	GGTTTCACCA	TGTTGGCCAG	GCTGGTTTTG	AACTCCTGAC
5851	CTCAGGTGAT	CCACCCGCCT	CAGCCTCCCA	AAGTGCTGAG	ATTATAGGTG
5901	TGAGCCACCA	CACCTGGCCT	CAGGAAGTAT	TTTTATTTTT	AAATTTATTT
5951	ATTTATTTGA	GATGGAGTCT	TGCTCTGTCG	CCCAGGCTAG	AGTGCAGCGA
6001	CGGGATCTCG	GCTCACTGCA	AGCTCCGCCC	CCCAGGTTCA	AGCCATTCTC
6051	CTGCCTCAGC	CTCCCGAGTA	GCTGGGACTA	CAGGCGCCCG	CCACCACACC
6101	CGGCTAATTT	TTTTGTATTT	TTAGTAGAGA	CGGGTTTTCA	CCGTGTTAGC
6151	CAGGAGGGTC	TTGATCTCCT	GACCTCGTGA	TCTGCCTGCC	TCGGCCTCCC
62.01	AAAGTGCTGG	GATTACAGGT	GTGAGCCACC	ACACCCGGCT	ATTTTTATTT
6251	TTTTGAGACA	GGGACTCACT	CTGTCACCTG	GGCTGCAGTG	CAGTGGTACA
6301	CCATAGCTCA	CTGCAGCCTC	GAACTCCTGA	GCTCAAGTGA	TCCTCCCACC
6351	TCATCCTCAC	AAGTAATTGG	GACTACAGGT	GCACCCCACC	ATGCCCACCT
6401	AATTTATTTA	TTTATTTATT	TATTTATTTT	CATAGAGATG	AGGGTTCCCT
6451	GTGTTGTCCA	GGCTGGTCTT	GAACTCCTGA	GCTCACGGGA	TCCTTTTGCC
6501	TGGGCCTCCC	AAAGTGCTGA	GATTACAGGC	ATGAGCCACC	GTGCCCAGCT
6551	AGGAATCATT	TTTAAAGCCC	CTAGGATGTC	TGTGTGATTT	TAAAGCTCCT
6601	GGAGTGTGGC	CGGTATAAGT	ATATACCGGT	ATAAGTAAAT	CCCACATTTT
6651	CTGGGCTTTA	TTTATTTATT	TATTTATTTA	TTTATTTTTA	ATTTTTTT
6701				TGGAGTGCAG	
6751				TCAAGCGATT	
6801				GCACCACCAT	
6851					AAGCTGGTCT
6901	· ·				GAGTGCTGGG
	•				



FIG. 10D

6951	ATTACAGGCA	TGAGCCACCA	TGCGTGGTCT	TTTTAAAATT	TTTTGATTTT
7001	TTTTTTTTT	GAGACAGAGC	CTTGCTCTGT	CGCCCAGGCT	GGAGTGCAGT
7051	GGCACGATCT	CAGCTCACTA	CAAGCTCCGC	CTCCCGGGTT	CACGCCATTC
7101	TTCTGCCTCA	GCCTCCTGAG	TAGCTGGGAC	TACAGGTGCC	CACCACCACG
7151	CCTGGCTAAT	TTTTTTTGGT	ATTTTTATTA	GAGACAAGGT	TTCATCATGT
7201	TGGCCAGGCT	GGTCTCAAAC	TCCTGACCTC	AAGTGATCTG	CCTGCCTCGG
7251	CCTCCCAAAG	CGCTGAGATT	ACAGGTGTGA	TCTACTGCGC	CAGGCCTGGG
7301	CGTCATATAT	TCTTATTTGC	TAAGTCTGGC	AGCCCCACAC	AGAATAAGTA
7351	CTGGGGGATT	CCATATCCTT	GTAGCAAAGC	CCTGGGTGGA	GAGTCAGGAG
7401	ATGTTGTAGT	TCTGTCTCTG	CCACTTGCAG	ACTTTGAGTT	TAAGCCAGTC
7451	GTGCTCATGC	TTTCCTTGCT	AAATAGAGGT	TAGACCCCCT	ATCCCATGGT
7501	TTCTCAGGTT	GCTTTTCAGC	TTGAAAATTG	TATTCCTTTG	TAGAGATCAG
7551	CGTAAAATAA	TTCTGTCCTT	ATATGTGGCT	TTATTTTAAT	TTGAGACAGA
7601	GTGTCACTCA	GTCGCCCAGG	CTGGAGTGTG	GTGGTGCGAT	CTTGGCTCAC
7651				TCTCGTGCCT	
7701		•		i .	ACTTTTGTAT
7751		GACAGGGTTT		GCTAAGCTGG	TCTCGAACTC
7801	CTGGCCTCAA			TCCCAAAGTG	CTGGGATTAC
7851	AGGTGTGAAC			TAGTGGCTTT	TAAGTGCTAA
7901		•			GTGAAGCATG
7951					GCAGCGTGGG
8001	AACTGGAAAG	-			GGGTCAGCCA
8051	•		AGACCATCTC		TGGGTCAGAG
8101				TCACAACCTT	
8151	AAAAAGCATT			AAAAAAGGCC	
8201	GTATGGATTT	· ·			GTTTTAGACA
8251	AGGAGCAGCT			•	GAGGAGCAGA
8301	GTGGATGTGG		•	GTCCCAGAGC	
8351	ACAGAGGGTT			AACTGAAATC	
8401	CAGTTCCAGA	•		ACGCTGCGGG	
8451	CTAGGGTTAC			GTAGGGGGAG	
8501	TGTTGGAAGA	•		TTTCAGGGCC	
8551				GGCCGAGGCG	
8601				ACATGGTGAA	
8651				GGTGGCGGGC	
8701				TGGCGTGAAC	· ·
8751				GCACTCCAGC	
8801				AAAGTAGGCT	
8851	GTGAGCTGAA				
8901	GACCCCTCGG				
8951	AGGGGAGACA		· · · · · ·		,
9001	TGGCTTTTAT				
9051	GAATACAGAA				
9101	•				TACTCTGTCA
9151	CCCAGGCTGA			· · · · · · · · · · · · · · · · · · ·	The state of the s
9201	CTGGGCTCAA				
9251	AGGTGCGTGT				** **
9301					GGATGCCCCT
	-010410010	COGNITICA	AUALCCCAUA	TYTOMIOGÓW	



FIG. 10E

9351	GTCTGCTGCC	TTGCCAGGGT	GCCAGGAGGG	CGCTGCTGTG	GAAGCTGAGG
9401	CCCGGCCATC	CAGGGCGATG	CATTGGGCGC	TGATTCTTGT	TCCTGCTGCT
9451	GCCTCGGTGC	TTAGCTTTTG	AAACAATGAA	ATAAATTAGA	ACCAGTGTGA
9501	AAATCGATCA	GGGAATAAAT	TTAATGTGGA	AATAAACTGA	ACAACTTAGT
9551	TCTTCATAAG	AGTTTACTTG	GTAAATACTT	GTGATGAGGA	CAAAACGAAG
9601	CACTAGAAGG	AGAGGCGAGT	TGTAGACCTG	GGTGGCAGGA	GTGTTTTGTT
9651	TGTTTTCTTT	GGCAGGGTCT	TGCTCTGTTG	CTCAGGCTGG	AGTACAGTGG
9701	CACAATCACA	GCTCACTATA	GCCTCGACCT	CCTGGACTCA	AGCAATCCTC
9751.	CTGCCTCAGC	CTCCCAGTAG	CTGGGACTAC	AGGCGCATGC	CACCATGCCT
9801	GGCTAATTTT	AAATTTTTTT	TTTTCTCTTT	TTTGAGATGG	AATCTCACTC
9851	TGTCGCCCAG	GCTGGAGTGC	AGTGGCGTGA	TCTCGGCTGA	CGGCAAGCTC
9901	CGCCTCCCAG	GTTCACTCCA	TTCGCCTGCC	TCAGCCTCCC	AAGTAGCTGG
9951	GACTACAGGC	GCTGGGATTA	CAAACCCAAA	CCCAAAGTGC	TGGGATTACA
10001	GGCGTGAGCC	ACTGCACCCG	GCCTGTTTTG	TCTTTCAATA	GCAAGAGTTG
10151	TGTTTGCTTC	GCCCCTACCT	TTAGTGGAAA	AATGTATAAA	ATGGAGATAT
10201	TGACCTCCAC	ATTGGGGTGG	TTAAATTATA	GCATGTATGC	AAAGGAGCTT
10251	CGCTAATTTA	AGGCTTTTTT	GAAAGAGAAG	AAACTGAATA	ATCCATGTGT
10301	GTATATATAT	TTTAAAAGCC	ATGGTCATCT	TTCCATATCA	GTAAAGCTGA
10351	GGCTCCCTGG	GACTGCAGAG	TTGTCCATCA	CAGTCCATTA	TAAGTGCGCT
10401	GCTGGGCCAG	GTGCAGTGGC	TTGTGCCTGA	ATCCCAGCAC	TTTGGGAGGC
10451	CAAGGCAGGA	GGATTCATTG	AGCCCAGGAG	TTTTGAGGCG	AGCCTGGGCA
10501	ATGTGGCCAG	ACCTCATCTC	TTCAAAAAAT	ACACAAAAAA	TTAGCCAGGC
10551	ATGGTGGCAC	GTGCCTGTAG	TCTCAGCTAC	TCAGGAGGCT	GAGGTGGGAG
10601		•		CAGTAAGCCA	TGATCTTGCC
10651	ACTGCATTCC	AGCCTGGATG	ACAGAGCGAG	ACCCTGTCTC	TAAAAAAAA
10701	AAAAACCAAA	CGGTGCACTG	TTTTCTTTTT	TCTTATCAAT	TTATTATTTT
10751	TAAATTAAAT	TTTCTTTTAA	TAATTTATAA	ATTATAAATT	TATATTAAAA
10801	AATGACAAAT	TTTTATTACT	TATACATGAG	GTAAAACTTA	GGATATATAA
10851	AGTACATATT	GAAAAGTAAT	TTTTTGGCTG	GCACAGTGGC	TCACACCTGT
10901	AATCCCAGCA	CTTTGGGAGG	CCGTGGCGGG	CAGATCACAT	GAGATCATGA
10951	GTTCGAGACC	AACCTGACCA	ACATGGAGAG	ACCCCATCTC	TACTAAAAAT
11001	ACAAAATTAG	CCGGGGTGGT	GGCGCATGCC	TGTAATCCCA	GCTACTCGGG
11051	AGGCTGAGGC	AGGAGAATCT	CTTGAACCCG	GGAGGCAGAG	GTTGCGGTGA
11101	GCCAAGATCG	TGCCTTTGCA	CACCAGCCTA	GGCAACAAGA	GCGAAAGTCC
11151	GTCTCAAAAA	AAAAGTAATT	TTTTTTAAGT	TAACCTCTGT	CAGCAAACAA
11201	ATTTAACCCA	ATAAAGGTCT	TTGTTTTTTA	ATGTAGTAGA	GGAGTTAGGG
11251	TTTATAAAAA	ATATGGTAGG	GAAGGGGGTC	CCTGGATTTG	CTAATGTGAT
11301	TGTCATTTGC	CCCTTAGGAG	AGAGCTCTGT	TAGCAGAATG	AAAAAATTGG
11351	AAGCCAGATT	CAGGGAGGGA	CTGGAAGCAA	AAGAATTTCT	GTTCGAGGAA
11401	GAGCCTGATG	TTTGCCAGGG	TCTGTTTAAC	TGGACATGAA	GAGGAAGGCT
11451	CTGGACTTTC	CTCCAGGAGT	TTCAGGAGAA	AGGTAGGGCA	GTGGTTAAGA
11501				CTAGACTAGC	
11551				TACTTTGAGT	
11601	•			TAGAGATATG	
11651				AATGGAGGAA	•
11701		•		AGTGTTTCCT	
11751				TGAGCTGAAT	
11801				AGTGCCCAGT	

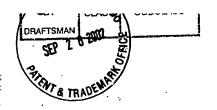


FIG. 10F

	•				
11851	GCTCTCTCAG	TGTTCCCTGA	TTGTTTTTTC	CTTTGTCATC	TTATCTACAG
11901	GATGTGACTG	GGAAGCTCTG	GTTTCAGTGT	CATGTGTCTA	TTCTTTATTT
11951	CCAGGCAAAG	GAAACCAACA	ATAAGAAGAA	AGAATTTGAG	GAAACTGCGA
12001	AGAAAGTGCG	CCGTGCCATC	GAGCAGCTGG	CTGCCATGGA	TTGAGGCCTC
12051	TGGCCGGAGC	TGCCTGGTCC	CAGAGTGGCT	GCACCACTTC	CAGGGTTTAT
12101	TCCCTGGTGC	CACCAGCCTT	CCTGTGGGCC	CCTTAGCAAT	GTCTTAGGAA
12151	AGGAGATCAA	CATTTTCAAA	TTAGATGTTT	CAACTGTGCT	CCTGTTTTGT
12201	CTTGAAAGTG	GCACCAGAGG	TGCTTCTGCC	TGTGCAGCGG	GTGCTGCTGG
12251	TAACAGTGGC	TGCTTCTCTC	TCTCTCTCTC	TTTTTTGGGG	GCTCATTTTT
12301	GCTGTTTTGA	TTCCCGGGCT	TACCAGGTGA	GAAGTGAGGG	AGGAAGAAGG
12351	CAGTGTCCCT	TTTGCTAGAG	CTGACAGCTT	TGTTCGCGTG	GGCAGAGCCT
12401	TCCACAGTGA	ATGTGTCTGG	ACCTCATGTT	GTTGAGGCTG	TCACAGTCCT
12451	GAGTGTGGAC	TTGGCAGGTG	CCTGTTGAAT	CTGAGCTGCA	GGTTCCTTAT
12501	CTGTCACACC	TGTGCCTCCT	CAGAGGACAG	TTTTTTTTTTT	GTTGTGTTTT
12551	TTTGTTTTTT	TTTTTTGGTA	GATGCATGAC	TTGTGTGTGA	TGAGAGAATG
12601	GAGACAGAGT			AACAACATGG	
12651	TTGTTTGAAT			CAAACTACAA	
12701	GCACAAAGCC	ATTCTAAGTC	ATTGGGGAAA	CGGGGTGAAC	TTCAGGTGGA
12751				TCTGGCAGAT	•
12801			•	AGCCGCGGGG	
12851				TAAATCCTTT	•
12901	•			TGCTGCAGGC	
12951				CCACACGGGG	
13001	GTCCGCCCAG			CAGCAGCTCC	•
13051	AAGTCTGGCG			GCCCTCCTCC	
13101	GCTGCAGGGT			GAAACCTCTG	
13151	CGGCTGTTCC			ATTTCAAACA	
13201				TTTCATCGTC	
13251	TGCCAACAGC			GTGAGGATGA	
13301	AGAGACGCAG			GAGCCACGAA	
13351	•			AAACAACTGA	•
13401				CGGCCTGAGA	
13451				TTTGTTAGCA	
13501				TTTTTGATAT	
13551	_		•	TTGCTTAGTA	
13601				TCAACTCCTT	
13651				CTGTGCTGTG	·
13701	•		•	CACGGCCTTT	
13751					AGCTTCCTCG
13801				CTAAGCGCAA	•
13851	•			TTTTCAGGTT	
13901				GCTTCTGGGC	
13951				CAACTGCCAT	
14001					CAAACTGGAG
14051		•			CAATTGCCAA
14101					CTTTTTTTT
14151			:		TGACCACAGC
14201					ACTTTAGCCT

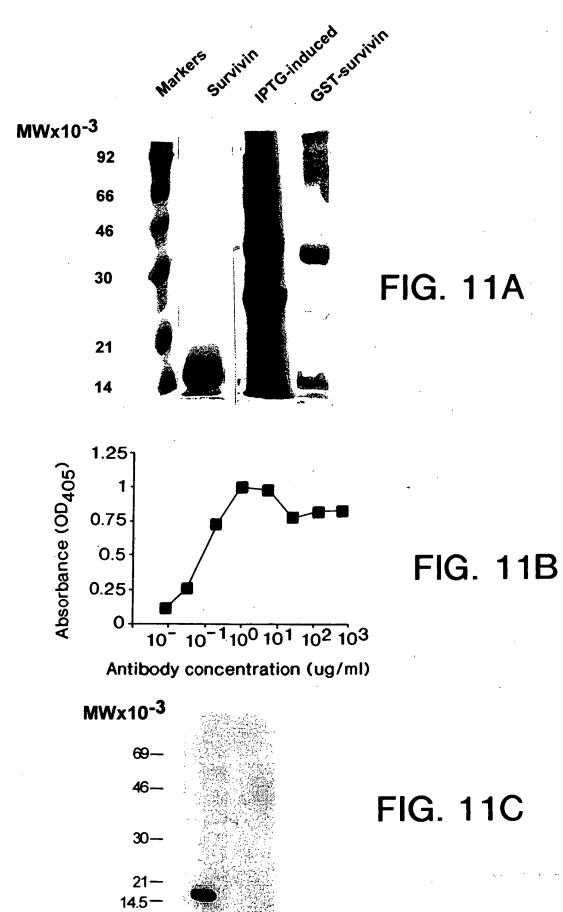


FIG. 10G

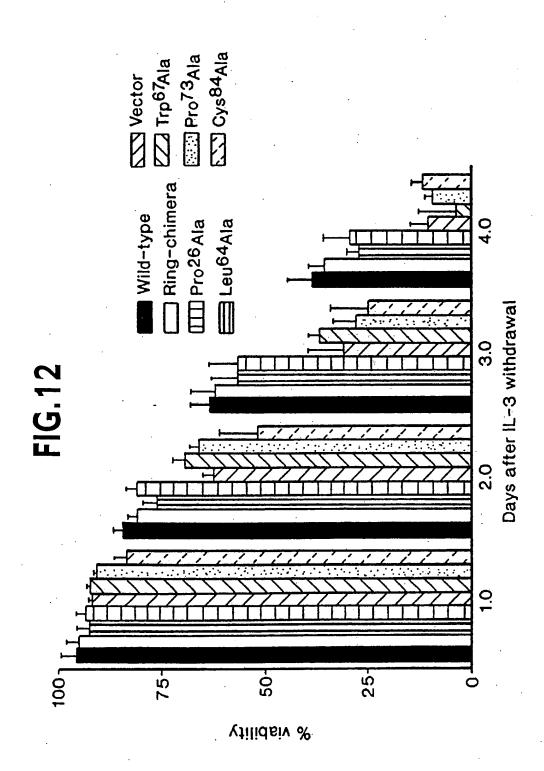
14251	CCCAAGTAGC	TGGGACTACA	GGCGCACGCC	ATCACACCCG	GCTAATTGAA
14301				ACTTTGTTGC	
14351				GCTTCAGCGT	
14401				CCTAGCTACC	
14451				ACTTGTCCAG	
14501	TAAGTAACTT	TTAGAGCTGG	GATTTGAACC	CAGGCAATCT	GGCTCCAGAG
14551				GCTTGGGAGG	
14601				GTCTTTAAAG	
14651	CATTTTCAGA	CAGCTTTGTC	AGAAAGGCTG	TCATATGGAG	CTGACACCTG
14701	CCTCCCCAAG	GCTTCCATAG	ATCCTCTCTG	TACATTGTAA	CCTTTTATTT
14751				CTAGTACAGG	

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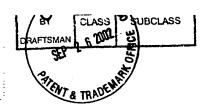












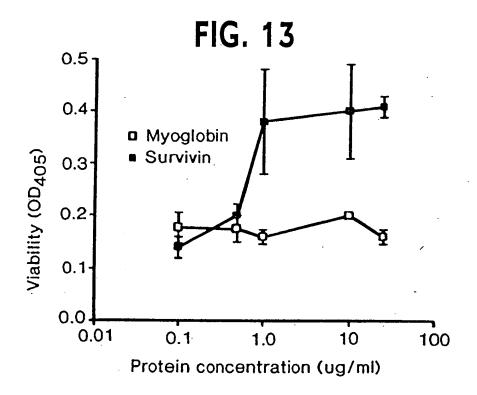
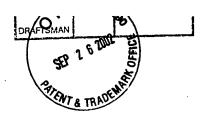
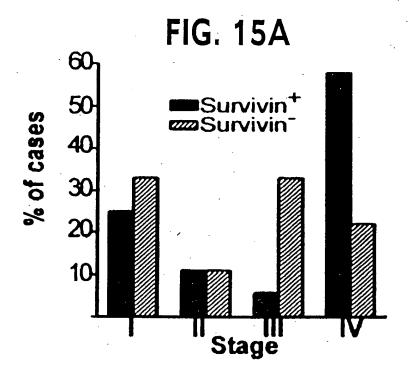


FIG. 14A FIG. 14B Prognostic factors Histology (Shimada class.) 757 75 n=72 % of survivin positive cases Survivin 50 % of cases 50 Survivin p=0.0025 25 Favorable Unfavorable 0 1 2 3 Number of negative prognostic factors (age, n-myc amplification Containing at least one widespread disease) negative prognostic factor





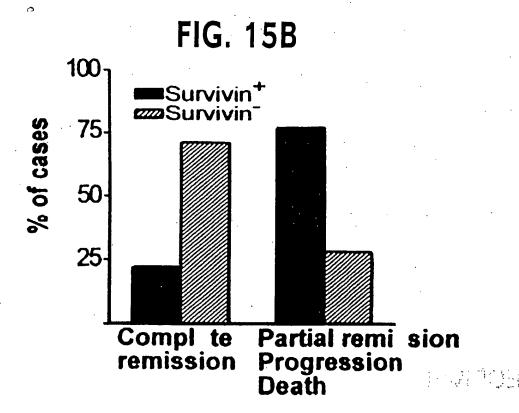
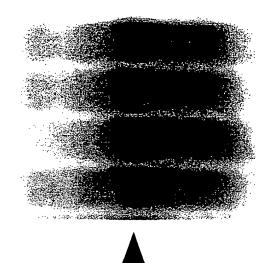
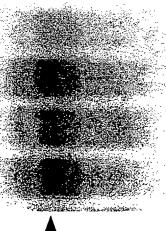


FIG. 16

TNHX DHX TNHY DHX





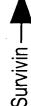




FIG. 17

